

AUTHOR'S CORRECTION

Multivariate Analyses Revealed Distinctive Features Differentiating Human and Cattle Isolates of Shiga Toxin-Producing *Escherichia coli* O157 in Japan

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Volume 49, no. 4, p. 1495–1500, 2011. Page 1495, abstract, line 12: “LSPA6 lineage I/II with *stx*₂ or *stx*₂ *stx*_{2c} strains” should read “LSPA6 lineage I/II with *stx*_{2c} or *stx*₂ *stx*_{2c} strains.”

Page 1495, abstract, line 13: “*stx*_{2c} or *stx*₂ *stx*_{2c} strains” should read “*stx*_{2c} or *stx*₁ *stx*_{2c} strains.”

Page 1497, left column, line 2: “except one isolate of LII” should read “except three isolates of LII.”

Page 1497, left column, line 6: “LII (36.7%)” should read “LII (36.4%).”

Page 1497, Table 2, third column (headed “Human”): The entry corresponding to LSPA6 genotype 222212 should read “2 (2.6),” the entry corresponding to “Subtotal” should read “3 (3.8),” the entry corresponding to LSPA6 genotype 231111 should read “3 (3.8),” and the entry corresponding to “Total” should read “78 (100).”

Page 1497, Table 2, fourth column (headed “Cattle”): The entry corresponding to “Total” should read “66 (100).”

Page 1497, Table 2, seventh column (headed “*stx*_{2c}”): The entry corresponding to “Subtotal” should read “15 (39.5).”

Page 1497, Table 2, eighth column (headed “*stx*₁ *stx*₂”): The entry corresponding to “Total” should read “61 (100).”

Page 1497, Table 2, ninth column (headed “*stx*₁ *stx*_{2c}”): The entry corresponding to “Subtotal” should read “12 (85.7).”